## **CLAIMS**

What is claimed is:

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1. An isolated nucleic acid comprising a nucleotide sequence encoding a polypeptide comprising an LRRCT domain consisting of the amino acid sequence:

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$$X_{12} C X_{13} X_{14} P X_{15} X_{16} X_{17} X_{18} X_{19} X_{20} D L X_{21} X_{22} L X_{23} X_{24} X_{25} D$$

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wherein X is any amino acid or a gap and the polypeptide does not comprise the amino acid sequence from residue 260 to 309 of SEQ ID NO: 5 (human NgR1) or SEQ ID NO: 17 (mouse NgR1).

- The isolated nucleic acid according to claim 1, wherein X<sub>17</sub> and X<sub>23</sub>
  are independently selected from the group consisting of: arginine and lysine.
  - 3. The isolated nucleic acid according to claim 2, wherein the amino acid sequence of the LRRCT domain is selected from the group consisting of: residues #261–310 of SEQ ID NO:2 and residues 261-310 of SEQ ID NO: 2 with up to 10 conservative amino acid substitutions.
    - 4. An isolated nucleic acid encoding the polypeptide of SEQ ID NO: 2.
- 5. An isolated nucleic acid encoding the polypeptide of SEQ ID NO: 4 (mouse NgR3) or SEQ ID NO: 14 (human NgR3).
  - 6. The isolated nucleic acid according to claim 1, wherein the

polypeptide comprises: (a) a NTLRRCT domain, and (b) less than a complete CTS domain, provided that a partial CTS domain, if present, consists of no more than the first 39 amino acids of the CTS domain.

- 7. The isolated nucleic acid to claim 1, wherein the polypeptide does not comprise an intact GPI domain.
- 8. An isolated nucleic acid consisting essentially of a nucleotide sequence complementary to a nucleotide sequence encoding a polypeptide selected from the group consisting of: a polypeptide consisting of residues 311-395 of SEQ ID NO: 2, a polypeptide consisting of residues 256-396 of SEQ ID NO:14 and a polypeptide consisting of residues 321-438 of SEQ ID NO: 4, wherein the nucleic acid is from 8 to 100 nucleotides in length.
  - 9. A vector comprising the nucleic acid of any one of claims 1, 4 or 5.
  - 10. A host cell comprising a vector according to claim 9.
  - 11. A polypeptide comprising a LRRCT amino acid sequence:

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$$N\; X_1\; W\; X_2\; C\; X_3\; C\; R\; A\; R\; X_4\; L\; W\; X_5\; W\; X_6\; X_7\; X_8\; X_9\; R\; X_{10}\; S\; S\; S\; X_{11}\; V$$

$$X_{12} C X_{13} X_{14} P X_{15} X_{16} X_{17} X_{18} X_{19} X_{20} D L X_{21} X_{22} L X_{23} X_{24} X_{25} D$$

25  $X_{26} X_{27} X_{28} C [SEQ ID NO: 19]$ 

wherein X is any amino acid residue or a gap and the polypeptide does not comprise the amino acid sequence from residue 260 to 309 of SEQ ID NO: 5 (human NgR1) or SEQ ID NO: 17 (mouse NgR1).

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12. The polypeptide according to claim 11, wherein  $X_{17}$  and  $X_{23}$  is selected from the group consisting of arginine and lysine.

- 13. The polypeptide according to claim 11, wherein  $X_{19}$  is glycine. [SEQ ID NO:11]
- 14. The polypeptide according to claim 11, wherein the amino acid sequence is selected from the group consisting of residues 261–310 of SEQ ID NO:2, residues 206–255 of SEQ ID NO: 14, residues 271–320 of SEQ ID NO:4 and amino acid sequences thereof comprising a conservative substitution.
  - 15. A polypeptide comprising a NTLRRCT amino acid sequence:

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- $C\ P\ X_{1}\ X_{2}\ C\ X_{3}\ C\ Y\ X_{4}\ X_{5}\ P\ X_{6}\ X_{7}\ T\ X_{8}\ S\ C\ X_{9}\ X_{10}\ X_{11}\ X_{12}\ X_{13}\ X_{14}\ X_{15}\ X_{16}\ P$   $X_{17}\ X_{18}\ X_{19}\ P\ X_{20}\ X_{21}\ X_{22}\ X_{23}\ R\ X_{24}\ F\ L\ X_{25}\ X_{26}\ N\ X_{27}\ I\ X_{28}\ X_{29}\ X_{30}\ X_{31}\ X_{32}\ X_{33}$   $X_{34}\ F\ X_{35}\ X_{36}\ X_{37}\ X_{38}\ X_{39}\ X_{40}\ X_{41}\ X_{42}\ L\ W\ X_{43}\ X_{44}\ S\ N\ X_{45}\ X_{46}\ X_{47}\ X_{48}\ I\ X_{49}$   $X_{50}\ X_{51}\ X_{52}\ F\ X_{53}\ X_{54}\ X_{55}\ X_{56}\ X_{57}\ L\ E\ X_{58}\ L\ D\ L\ X_{59}\ D\ N\ X_{60}\ X_{61}\ L\ X_{62}\ X_{63}\ X_{64}$   $X_{65}\ P\ X_{66}\ T\ F\ X_{67}\ G\ L\ X_{68}\ X_{69}\ L\ X_{70}\ X_{71}\ L\ X_{72}\ L\ X_{73}\ X_{74}\ C\ X_{75}\ L\ X_{76}\ X_{77}\ L\ X_{78}$   $X_{79}\ X_{80}\ X_{81}\ F\ X_{82}\ G\ L\ X_{83}\ X_{84}\ L\ Q\ Y\ L\ Y\ L\ Q\ X_{85}\ N\ X_{86}\ X_{87}\ X_{88}\ X_{89}\ L\ X_{90}\ D\ X_{91}\ X_{92}\ F\ X_{93}\ D\ L\ X_{94}\ N\ L\ X_{95}\ H\ L\ F\ L\ H\ G\ N\ X_{96}\ X_{97}\ X_{98}\ X_{99}\ X_{100}\ X_{101}\ X_{102}$   $X_{103}\ X_{104}\ F\ R\ G\ L\ X_{105}\ X_{106}\ L\ D\ R\ L\ L\ L\ H\ X_{107}\ N\ X_{108}\ X_{109}\ X_{110}\ X_{111}\ V\ H\ X_{112}$   $X_{113}\ A\ F\ X_{114}\ X_{115}\ L\ X_{116}\ R\ L\ X_{117}\ X_{118}\ L\ X_{119}\ L\ F\ X_{120}\ N\ X_{121}\ L\ X_{122}\ X_{123}\ L$   $X_{124}\ X_{125}\ X_{126}\ X_{127}\ L\ X_{128}\ X_{129}\ L\ X_{130}\ X_{131}\ L\ X_{132}\ X_{133}\ L\ R\ L\ N\ X_{134}\ N\ X_{135}\ W$   $X_{136}\ C\ X_{137}\ C\ R\ X_{138}\ R\ X_{139}\ L\ W\ X_{140}\ W\ X_{141}\ X_{142}\ X_{143}\ X_{144}\ R\ X_{145}\ S\ S\ S\ X_{146}$   $V\ X_{147}\ C\ X_{148}\ X_{149}\ P\ X_{150}\ X_{151}\ X_{152}\ X_{153}\ X_{154}\ X_{155}\ D\ L\ X_{156}\ X_{157}\ L\ X_{158}\ X_{159}\ X_{160}$   $D\ X_{161}\ X_{162}\ X_{163}\ C\ [SEQ\ ID\ NO:18]$
- wherein X is any amino acid residue or a gap and wherein the polypeptide is not the polypeptide of SEQ ID NO: 5 (human NgR1) or SEQ ID NO: 17 (mouse NgR1).
- 16. The polypeptide according to claim 15, wherein  $X_6$ ,  $X_{37}$  and  $X_{38}$  30 represents a gap.
  - 17. A polypeptide comprising an amino sequence selected from the

group consisting of: SEQ ID NO:2, SEQ ID NO:4 and SEQ ID NO:14.

- 18. The polypeptide according any one of claims 11, 15 or 17, wherein the polypeptide comprises: (a) an NTLRRCT domain, and (b) less than a complete CTS domain, provided that a partial CTS domain, if present, consists of no more than the first 39 amino acids of the CTS domain.
  - 19. The polypeptide according to any one of claims 11, 15 or 17, wherein the polypeptide does not comprise an intact GPI domain.

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- 20. The polypeptide according to any one of claims 11, 15 or 17, wherein the amino acid sequence of the polypeptide further comprises an amino acid sequence of a heterologous polypeptide.
- 15 21. The polypeptide according to claim 20, wherein the heterologous polypeptide is an Fc portion of an antibody.
- 22. A method of producing a polypeptide according to any one of claims 11, 15 or 17, comprising the steps of introducing an isolated nucleic acid according to any one of claims 1, 4, 5 or 8 or a vector according to claim 9 into a host cell, culturing said host cell under conditions suitable for expression of said polypeptide, and recovering said polypeptide.
- 23. An antibody that binds to a polypeptide of any one of claims 11, 15 or 17.
  - 24. A composition comprising the polypeptide of claim 11, 15 or 17 and a pharmaceutically acceptable carrier.
  - 25. A composition comprising the antibody of claim 23 and a pharmaceutically acceptable carrier.

- 26. A method of decreasing inhibition of axonal growth of a CNS neuron, comprising the step of contacting the neuron with an effective amount of the polypeptide of claim 11, 15 or 17.
- 5 27. A method of treating a central nervous system disease, disorder or injury, comprising administering to a mammal an effective amount of the polypeptide of claim 11, 15 or 17.
- 28. A method of decreasing inhibition of axonal growth of a CNS neuron comprising the step of contacting the neuron with an effective amount of the antibody according to claim 23.
- 29. A method of treating a central nervous system disease, disorder or injury, comprising administering to a mammal an effective amount of the antibody
  according to claim 23.
  - 30. A method for identifying a molecule that binds a polypeptide of claim 11, 15 or 17 comprising the steps of:
    - (a) providing a polypeptide of claim 11, 15 or 17;
    - (b) contacting the polypeptide with the candidate molecule;
      - (c) detecting binding of the candidate molecule to the polypeptide.

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